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 M E E D L F Q L R Q L P V 13
 AGGGCTCATCCTGGCGAGCGGGAC ATG GAG GAG GAC TTA TTC CAG CTA AGG CAG CTG CCG GTT 39
 V K F R R T G E S A R S E D D D T A S G E 33
 GTG AAA TTC CGT CGC ACA GGC GAG AGT GCA AGG TCA GAG GAC GAC ACG GCT TCA GGA GAG 39
 H E V Q I E G V H V G L E A V E L D D G 53
 CAT GAA GTC CAG ATT GAA GGG GTC AAC GTG GGC CTA GAG GCT GTG GAG CTG GAT GAT GAT GGG 159
 A A V P K E F A N P T D D T F M V E D A 73
 GCA GCT GTG CCC AAG GAG TTT GCC AAT CCC ACC GAT GAT ACT TTC ATG GTG GAA GAT GCA 219
 V E A I G F G K F Q W K L S V L T G L A 93
 GTG GAA GCC ATT CGC TTT CGA AAA TTT CAG TGG AAG CTG TCT GTT CTC ACT GGC TTG GCT 279
 W M A D A M E M M I L S I L A P Q L H C 113
 TGG ATG GCT GAT GCC ATG GAG ATG ATC CTC AGC ATC CTG GCA CCA CAG CTG CAT TGC 339
 E W R L P S W Q V A L L T S V V F V G M 133
 GAG TGG AGG CTC CCA AGC TGG CAG GTG GCA TTG CTG ACC TCG GTG GTC TTT GTA GGC ATG 399
 M S S S T L W G N I S D Q Y G R K T G L 153
 ATG TCC AGC TCC ACG CTC TGG GGA AAT ATC TCA GAC CAG TAC GGC AGG AAA ACA GGG CTG 459
 K I S V L W T L Y Y G I L S A F A P V Y 173
 AAG ATC AGC GTG CTG TGG ACT CTG TAC TAT GGC ATC CTT AGT GCA TTT GCG CCC GTG TAT 519
 S W I L V L R G L V G F G I G G V P Q S 193
 AGC TGG ATC CTG GTG CTC CGG GGC CTG GTG GGC TTC CGG ATC CGA GGA GTT CCC CAG TCG 579
 V T L Y A E F L P M K A R A K C I L L I 213
 GTG ACG CTG TAT GCC GAG TTC CCC ATG AAA GCC AGA GCT AAA TGT ATT TTG CTG ATT 639
 E V F W A I G T V F E V V L A V F V M ? 233
 GAG GTA TTC TGG GCC ATC GGG ACA GTG TTC GAG GTC GTC CTG GCT GTG TTC CTG ATG CCC 699
 S L G W R W L L I L S A V P L L L F A V 253
 AGC CTG GGC TGG CGT TGG CTG CTC ATC CTC TCA GCT GTC GTC CTC CTC CTC TTT GCC GTG 759
 L C F W L P E S A R Y D V L S G N Q E K 273
 CTG TGT TTC TGG CTG CCT GAA AGT GCA AGG TAT GAT GTG CTG TCA GGG AAC CAG GAA AAG 819
 A I A T L K R I A T E N G A P M P L G K 293
 GCA ATC GCC ACC TTA AAG AGG ATA GCA ACT GAA AAC CGA GCT CCC ATG CCG CTG GGG AAA 879
 L I I S R Q E D R G K M R D L F T P H F 313
 CTC ATC ATC TCC AGA CAG GAA GAC CGA GGC AAA ATG AGG GAC CTT TTC ACA CCC CAT TTT 939
 R W T T L L W F I W F S N A F S Y Y G 333
 AGA TGG ACA ACT TTG CTG CTG TGG TTT ATA TGG TTT TCC AAT GCA TTC TCT TAC TAC CGG 999

Figure 1

DRAFT

L V L I T T E S C F Q A G D V C G I S S R	353
TTA GTT CTA CTC ACC ACA GAA CTC TTC CAG GCA GGA GAT GTC TCC CCC ATC TCC AGT CGG	1059
K K A V E A K C S L A C E Y L S E E D Y	373
AAG AAG GCT GTC GAG GCA AAA TGC AGC CTG GCC TGC GAG TAC CTG AGT GAG GAG GAT TAC	1119
M D L L W T T L S E F P G V L V T L W I	393
ATG GAC TTG CTG TGG ACC ACC CTC TCT GAG TTT CCA GGT GTC CCT GTG ACT CTG TGG ATT	1179
I D R L G R K K T M A L C F V I F S F C	413
ATT GAC CGC CTG GGG CCC AAG AAG ACC ATG GCC CTG TGC TTT GTC ATC TTC TCC TTC TGC	1239
S L L L F I C V G R N V L T L L L F I A	433
AGC CTC CTG CTG TTT ATC TGT GGT GGA AGA AAT GTG CTC ACT CTG TTA CTC TTC ATT GCA	1299
R A F I S G G F Q A A Y V Y T P E V Y P	453
AGA GCG TTT ATT TCT GGA GGC TTT CAA GCG GCA TAT GTT TAC ACA CCT GAG GTC TAC CCC	1359
T A T R A L G L G T C S G M A R V G A L	473
ACG GCA ACG CGG GCC CTC CGC ACC TGC AGC GGC ATG GCA AGA GTG GGT GCT CTC	1419
I T P F I A Q V M L E S S V Y L T L A V	493
ATC ACT CCG TTC ATC GCC CGG GTG ATG CTG GAA TCC TCT GTG TAC CTG ACT CTG GCA GTT	1479
Y S G C C L L A A L A S C F L P I E T K	513
TAC AGT GGC TGC TGC CTC CTG GCT GCC CTG GCC TCC TGC TTT TTG CCC ATT GAG ACC AAA	1539
G G G L Q E S S H R E W G Q E M V G R G	533
GGC GGA GGA CTG CAG GAG TCC AGC CAC CGG GAG TGG GGC CAG GAG ATG GTC GGC CGA GGA	1599
M H G A G V T R S N S G S Q E *	549
ATG CAC GGT GCA GGT GTT ACC AGG TCG AAC TCT GGC TCT CAG GAA TAG	1647
TGACCGATGGGGACTGAGCTGGCTTGAGGCTCCAGAGCTGGGGGCTGGAGCCCCACTGGGCACTGATTGT CACTGCCACATCAAGAACCTACCCAGAGTATGACCTGGACCAACAGGGTTTGTGCTTGACTCAGTTGCTCATCT TCATTGAGGTCCACCCAGGGATGGGAGATGTTCTCTAGGGGTTCTCTATATGTGGTAAAGCTTGTCTAA CCTGTGGATCTACATGGGAAGACTACCCATATTAGGAGGGTCTGGTAATGCCAGCAACCAATCAGACACCAGAGT CACCCGCCAACCCCTCAGTAACAACAAAATATCTCTGTAGATAACCGTCCAGGCTCAGGCCATGTGACACCTGC TGTCCACCCACCGGACCTGTCAGTAGGTTCTCCCACACCCACAGCCCCAGGTTTCTCTTGTAAAGCTTGGGAT CTAGGTGTGGCTGAGCAGCTATTCTGGCAGGGGCCCCCGGTTTGCTCCCTAGAGCCTGACCAGTGGATTCTCTG GCAGATGGACATGGTGCATTCAAAGTGGAGCCACATGCCACCCAGGCCNTGGAGTTGCCGTTGGCACCA AGAGATCCAGATGTGTCCTGGGACAGCTGGGTCTTCCACCAGGTGACAACCTCAAAACGCCGTTACCCCTGGGAAC TGAGGACTGAGSGCCAAGTG	

Figure 1 continued

0 0 0 0 0 0 0 0 0 0 0 0 0

*wCFMAaFGGmFMFGWDTGVINGFvwMIdFhYRFgMmhydwtyYl....S
+ A+G F W V G +WM D + ++ ++ ++
132 EDAVEAIGF-GKFQWKLSVLTGLAWMADAMEMMILSILAPQLHCEWRLP 179

TMRWgLIVSIFnIGCMIGSIFFGWigDMYGRRmsMMMVnvIFIIGIIIMI
+ + +L +S ++G+M +S ++G+I+D YGR+ ++ ++++ +I+++
180 SWQVALLTSVVFVGMMSSSTLWGNISDQYGRKTGLKISVLWTLYYGILSA 229

fSinysWWMYIIGRIImGICVGgISvlvPMYISEIAPkH1RGtMvSWYQL
F++ + +++ R ++G+G+GG + V +Y +E+ P + R +++ + ++
230 FAP--VYSWILVLRGLVGFIGGVPQSVTLY-AEFLPMKARAKCILLIEV 276

MITiGIFIAYCfNYgfnYnNdsWQWRWPLGLcFIWAIfMIIgMMF1PES
+++IG ++ + +++ + + WRW+L+L++++ +++ +++++LPES
277 FWAIGTVFEVVLAVFVMPSLG---WRWLLILSAVPLLLFAVLCFWLPES 322

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R+ V+ G+ E+A ++L+R+ ++ + + + + + + + + + + + + + + + +
323 ARYDVLSGNQEKAIA TLKRIATEENGAPMPLGKLII SRQED-----RGK 365

WgELFr rr rtPkMRWRiIMgMMIQiFQQFTGINYIMYYsTTIFesVGMq*
+ LF+ + RW ++ + I+++ ++ YY+ ++ ++ + Q
366 MRDLFTP HF---RWTTLLLWFIWFSN-----AFSYYGLVLLTTELFQ 404

Figure 2

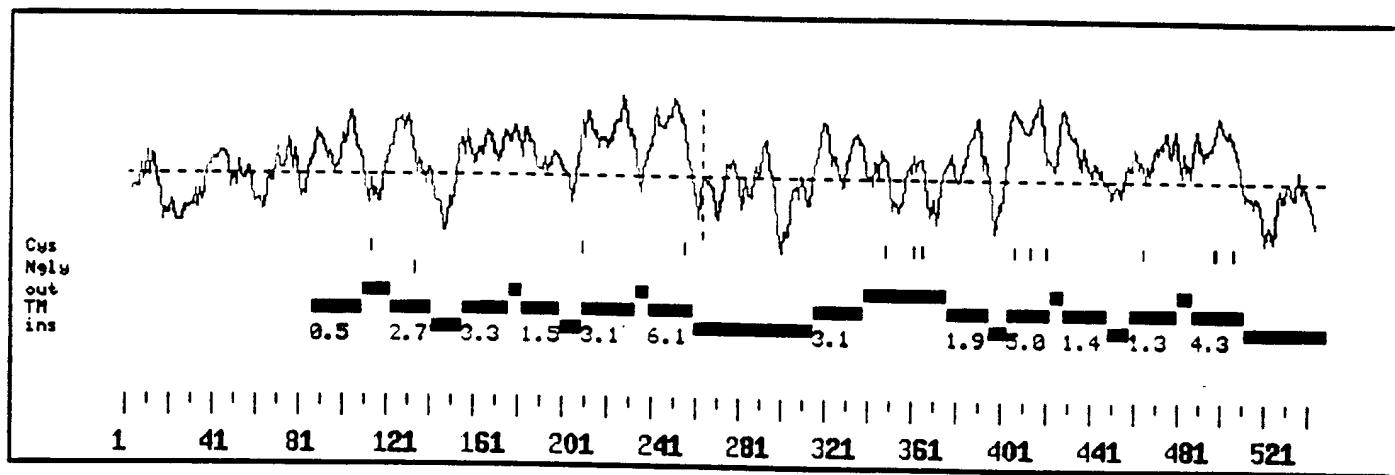


Figure 3

GTCGACCCACCGTCCGAGCAAAGAGGATTACATGGACCTGCTGTGGACCACCCGTCTG
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TGGCTCTGTGTTCGCATCTTTCCTCTGCAGCCTCTGCTGTTCATCTGCATTGGAA
GAAATGTGCTAACCCCTTACTGTTCATGCAAGAGCGTTTATTTCTGGAGGCTTCCAAG
CAGCCTACGTTACACGCCGTAGGGTGTATCCAACGGCGACGAGGGCGCTGGGCCTGGGCA
CCTGCAGCGGCATGGCGAGAGTGGGCGCTCATCACTCCATTCACTAGCTCAGGTGATGC
TGGAAATCTCCGTACCTGACCCCTGGCGTCTACAGTGGCTGCTGCCCTCTGCTGCCCT
TGGCCTCTGTTCTGCCCATCGAGACCAAAGGCCGAGCACTGCAGGAGTCCAGGCCACC
GGGAGTGGGGCCAGGAGATGGTGGCCAGGGACAAACAGCACAGCGTCCCAGGTGCA
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CAGGAGGCTGTTGTCTGGCCTCAGGAGGAAGGTAGGTCTGAAGGCAGTCCACGTGTACT
CCCGGATTCCGAAACGCCACGAGCCGCCCTCTGAGGATTAGGAAAGAAGCAGCAGGGGG
AGGAAGGAGATGGCCAGGCCAAATCAAAGGCCAGAGGAAGTGGCCGCTTGTGATG
GACACCTCGCAAGGGAGCCAAATGTGCGTTGTGCCCTCATCTCTTA

Figure 4